\_\_\_\_\_\_

Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: [year=2009; month=4; day=28; hr=13; min=38; sec=23; ms=829; ]

\_\_\_\_\_\_

\*\*\*\*\*\*\*\*\*\*\*\*\*

Reviewer Comments:

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Arg Gly Asp Ser

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The above is an insufficient response for numeric identifier <223>. Please explain the source of the genetic material. When using "Unknown" for numeric identifier <213>, please provide as much taxonomic information, as possible, about the organism from which the genetic material was extracted. If the genetic material was extracted from a sample in which there was an unknown variety of organisms, please explain where the sample was taken, for example a soil sample. These errors appear in other sequences in the sequence listing. Please make all necessary changes.

## Validated By CRFValidator v 1.0.3

Application No: 10593484 Version No: 1.0

Input Set:

Output Set:

**Started:** 2009-04-25 15:06:57.002 **Finished:** 2009-04-25 15:06:57.810

**Elapsed:** 0 hr(s) 0 min(s) 0 sec(s) 808 ms

Total Warnings: 14

Total Errors: 0

No. of SeqIDs Defined: 14

Actual SeqID Count: 14

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## SEQUENCE LISTING

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      Jochen KLOCK
      Philippe Emmanuel MAILLAN
       Juergen H. VOLLHARDT
       Raphael BEUMER
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